

F/G.1

Receptor Construct:	WT		D4D			
Antibody applied for Immunoprecipitation:	318	67	13	318	67	13
		***		•		-
				•	•	

ctg ago Leu ser

gċg

ťgċ Čvs

Gln. Val. Glu! Thr

cag.gtg.gaa.act

ggc.gtg.gcc.aga

ttċ

Phe

Gly .Val.Ala.Arg

## F/G.

14

agggcgcgag ggcgcgaggg cagggggcaa ccggaccccg agagaaggcg ctgggctgcg cggagcctgg gcgagcgcag

gga ctg g gtc tgg gcc gcg ctg gcc gtc Val Trp Ala Ala Leu Ala Val Ala Val gcc Val gtc Met Ala Pro S atg gcg cccgcaccc

gag ctc tgg gct gcg Glu Leu Trp Ala Ala gcc ttg ccc -22 Cac gcg 147

aca tgc cgg caa cat gca agc Ser ggc. ccg 999 61y gcc ccg gag ccc Ala Pro Glu Pro aaa tgc tcg agc. tac tgc tgc Pro Tyr cag atg t ttt aca Phe Thr gca Ala gct aca gtg Vai cag cag gcc Ala gac Pro tat His Ala Leu tac gaa ctc aga .279. Ala

Leu. Arg. Glu. Tyr. Asp. Gln. Thr. Ala. Gln. Met. Cys. Cys. Ser. Lys. Cys. Ser. Pro. Gly. Gln. His, Ala agc aca.tac.acc.cag tcc.tgt.gag.gac gac gac.acc.gtg tgt tcg acc.aag.acc tgt ttc aaa.gtc .Lys.Val

Ser. Thr. Tyr. Thr. Gln Ser.Cys.Glu.Asp Asp Asp. Thr. Val. Cys Ser Thr, Lys. Thr

54. Cys phe

.345.

tac fyr cgc:tgt:agc,tct gac Arg. Cys. Ser. Ser. Asp tgg ccc ggc 1 Pro Gly agg Ser . 9/ ŧgċ tcc Çys. acc 386 Leu.Ser.Cys.Gly Thr. àtc tgc ttg.agc.tgt ile. Arg çgç tgc Cys àaċ Àsń cag r'Glu'ch gtt.ccc.gag Val. Pro.Glu gaa Trp tgg Ťhř act tgċ Ċys aac A.sn .ctc.tgg Leu. Trp caa gcc .411.

Arg aag Cys .tgc 999.tgc.cgg.ctg gag Gln Ala 477

tgc.cgc.ccg.ggc Cys. Arg. Pro. Gly Z/S gcg.ccg.ctg.cgc ζys .Cys.Arg.Leu Gly Glu aag.cag Lys.Gln 543.

tto Phe .೦೮೮. ರೆವಿರೆ. ಇಂಡ Pro Gly Thr gcc Cys Ala 120 tgt ပ္ပပ္ Val.Cys:Lys.Pro gtg tgc.aag gtg Val aca.tca.gac Thr. Ser. Asp gaa Glu act Thr cca.gga Pro.Gly 609.

Pro gca Àla gtg gcc atc Val. Ala Ile 999 Pro. CCa Ala Ser Met Val Cys' Asn agt .164 <u>66</u>5 Thr Ile acc. .000 pro. Ġl'n. Hi ŝ tcć Ser acg tcc acg Thr Ser Thr Cys Arg Pro tgc Cys Ile gtc Asp gca  $\operatorname{Th} \dot{\Sigma}$ atg.gat. Asp Ser Ser Met

gca agc

ģġġ Glý

ćcċ

gtģ

tgt aac

a C.

ċaġ

Ċac

ccc.

tġc aġg.

att.

ġať

àcg

tca tcc

142.

Ser

Alà

Asn aat

cac tta His Leu

gta

Thr

Ser Asn Thr. ttc.aac.acg.

act

## F16.2B

TRANSMEMBRANE					
cag cca gtg tcc aca cga tcc caa cac acg cag cca act cca gaa ccc agc act gct cca agc acc acc gln Pro Val Ser Thr Arg Ser Gln His Thr Gln Pro Thr Pro Glu Pro Ser Thr Ala Pro Ser Thr Ala Pro Ser Thr Ala Pro Ser Thr Bro Glu Pro Ser Thr Gly Asp Rec at gcc cc agc ccc agc ccc agc gct gaa ggg agc act ggc gac act gcc gcc acc acc agc ccc agc ccc agc gcc acc gcc acc gcc acc gcc acc gcc acc a	acc cag gtg aaa aag aag ccc ttg tgc ctg cag aga gaa gcc aag gtg cct cac Thr Gln Val Lys Lys Lys Pro Leu Cys Leu Gln Arg Glu Ala Lys Val Pro His 274 aag gcc cgg ggt aca cag ggc ccc gag cag cag cac ctg ctg atc aca gcg ccg Lys Ala Arg Gly Thr Gln Gly Pro Glu Gln Gln His Leu Leu Il- Thr Ala Pro	ago ago too ctg gag ago tog goo agt gog ttg gac aga agg gog coc act cgg aac cag coa cag ser Ser Ser Leu Glu Ser Ser Ala Leu Asp Arg Arg Ala Pro Thr Arg Asn Gln Pro Gln 1137  318  900 coa ggo gtg gag gcc gcg gag gcc cgg gcc agc acc ggg ago toa gat tot toc Gly Ala Gly Glu Ala Arg Ala Ser Thr Gly Ser Ser Ser Pro Gly Gly His Gly Thr Gln 1203	ctt ggt ggc cat ggg acc cag gtc aat gtc acc tgc atc gtg aac gtc tgt agc agc tct gac cac Ala Pro Gly Val Glu Ala Ser Val Asn Val Thr Cys Ile Val Asn Val Cys Ser Ser Ser Asp His 1269 agc tca cag tgc tcc tcc caa gcc agc tcc aca atg gga gac aca gat tcc agc ccc tcg gag tcc Ser Ser Ser Gln Ala Ser Ser Thr Met Gly Asp Thr Asp Ser Ser Pro Ser Glu Ser 1335	gac gag cag gtc ccc ttc tcc aag gag gaa Asp Glu Gln Val Pro Phe Ser Lys Glu Glu Ctg ctg ggg agc acc gaa gag aag ccc ctg Leu Leu Gly Ser Thr Glu Glu Lys Pro Leu taa ccaggccggt gtgggctgtg tcgtagccaa ggt	Ser End 439

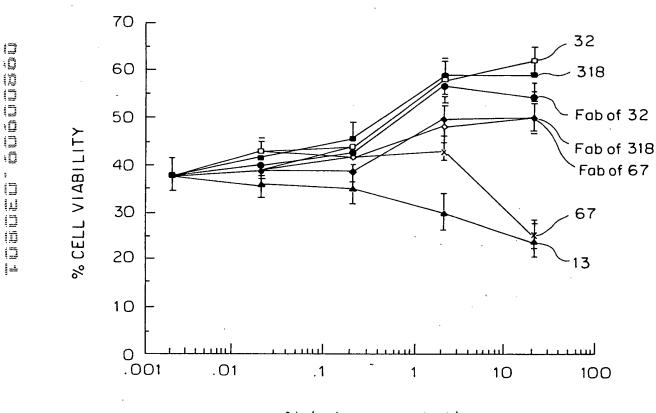
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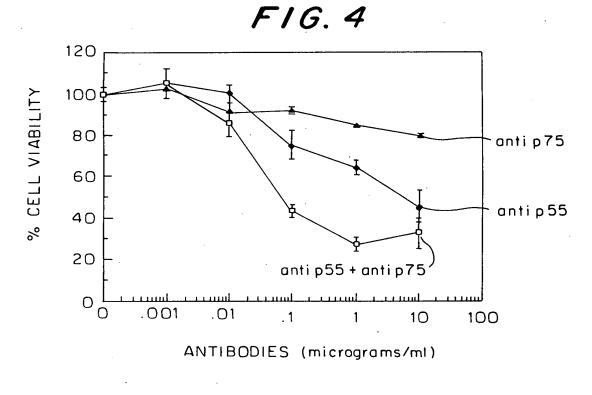
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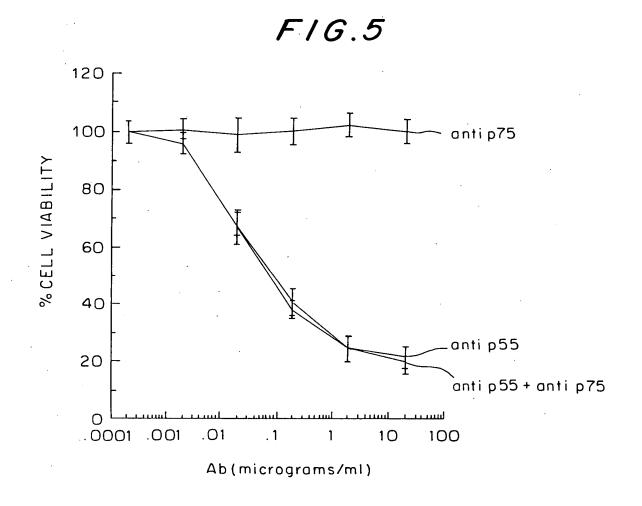
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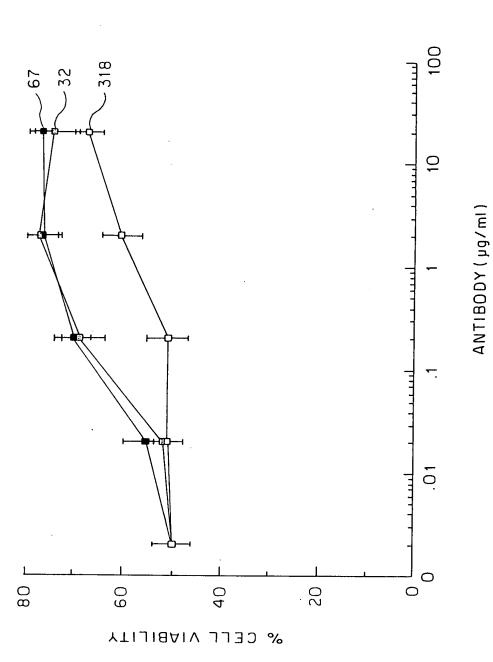
FIG.3

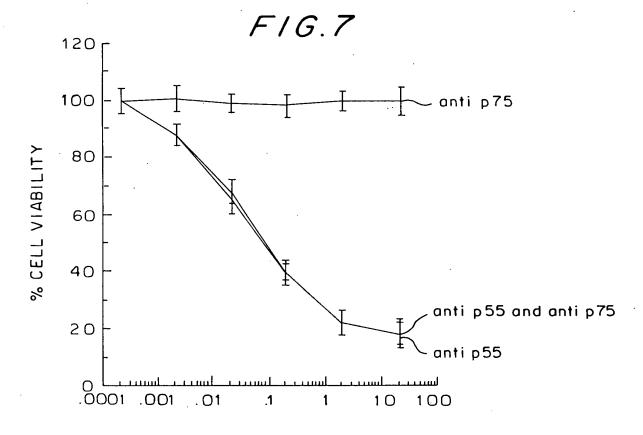


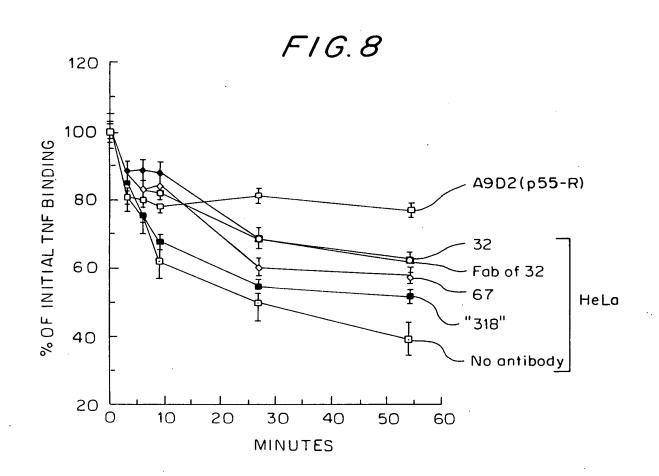




F16.6







Ab (micrograms/ml)

APPROVED O.G. FIG.

BY CLASS SUBCLASS

DRAFTSMAN

## F16.9

> 61 . 1 H	GERREO	181005	<u>  FBBB</u>
QNLEGLH-HDGQFCH-KPCPPGERKARDCTVNGDEPDCV ACPTGLYTHSGECC-KACNLGEGVAQPCGANOTVCE ACREKQYLINSQCC-SLCQPGQKLVSDCTEF-TETECL NCVKDTYPSGHKCC-RECQPGHGMVSRCDHT-RDTVCH	ECESGSFTASERHL-RHCLSC-SKCRKENGOVEISSCTVD-RDTVC SCEDSTYTQLWNWV-PECLSCGSRCSDD-QVETQACTRE-QNRIC PCQEGKEYTDKAHFSSKCRRC-RLCDEGHGLEVEINCTRT-QNTKC PCLDSVISSDVVSATEPCKPC-TECVGLQSHSAPCVEA-DDAVC PCGESEFLDIWHREIN-CHQH-KYCDPNLGLRVQQKGTSE-TDTIC PCGESFFLDIWHREIN-CHQH-KYCDPNLGLRVQQKGTSE-TDTIC	-CRKNQYRHYWSENLFQCFNCSLCLHGT-VHLSCQEK-QNTVCCRPGWYCALSKQEGCRLCAPLRKCRPGFGVARPGTET-SDVVCK-CKPPGFGVARPGTET-SDVVCK-CKPPGFGVARPGTET-SDVVCK-CKPPGFFGVARPGTET-SDVVCK-CKPPGFFGVARPGTLT-SNTKCCAYGYYQDETTGRCEACRVCEAGSGLVFSCQDK-QNTVCE-CEEGWHCTSEACESCVLHRSCSPGFGVKQIATGV-SDTICE-CRPGFTQPRQDSCV	TCHAGFFLRENECVSC-SNCKKSLECTKLC PCAPGTFSNTTSST-DICRPH-QICNVVAIPGNASMDAVC ECPDGTYSDEAHHV-DPCLPC-TVCEDTERQLRECTRW-ADAEC PCPVGFFSNVSSAF-EKCHPTSCETKDLVVQQAGTNKTDVVCC PCPPGHFSPGSHQACKPW-TNCTLSGKQIRHPASNSLDTVC
NGF-R (3-31) CDw40 (25-60) Ox40 (25-60)	p55 TNF-R(4.3-86) p75 TNF-R(77-119) FAS (68-112) NGF-R (38-80) CDw40 (61-104) Ox40 (61-104)	p55 TNF-R(87-126) p75 TNF-R(120-162) FAS (113-149) NGF-R (81-119) CDw40 (105-144) Ox40 (105-123)	p55 TNF-R(127-155) p75 TNF-R(163-201) NGF-R (120-161) CDw40 (145-186) Ox40 (124-164)
	hu hu hu rat	hu hu hu hu hu rat	hu hu hu hu hu hu rat